1/9

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GGATTC

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Fig. 1

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1 EEA S GKVVQD	IMPCLHFVKG	eekepske ss	SGTKKLSEEV	/ ~	
51 KCIVRATKGI	SGIKNELVAE	VPKKCDIKTT	LPPITADFDC	_	L00 3
101 YY <i>GF</i> QET <u>S</u> GT	MVRALMPCLP	FVQGKEKEPS	KG SS SGAKRL	_	150 R
151 VHACECIQTA	MKTYSDIDGK	LVSEVPKHCG	IVDSKLPPID		200 V
201 VPROPOLPVS	LRHGPVTGPS	DPAHKARLER	PQIRVPPPAP	243 EKA	

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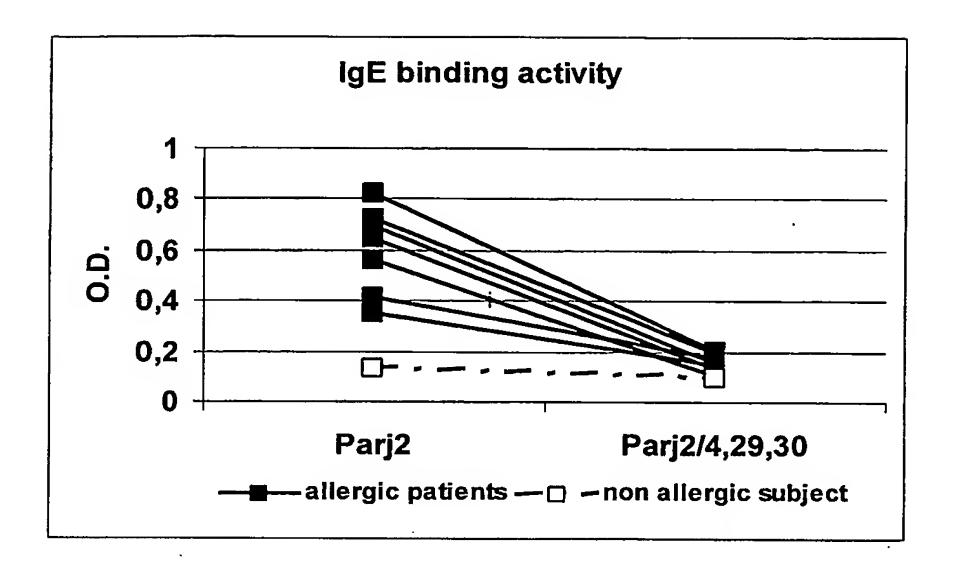


FIG. 3

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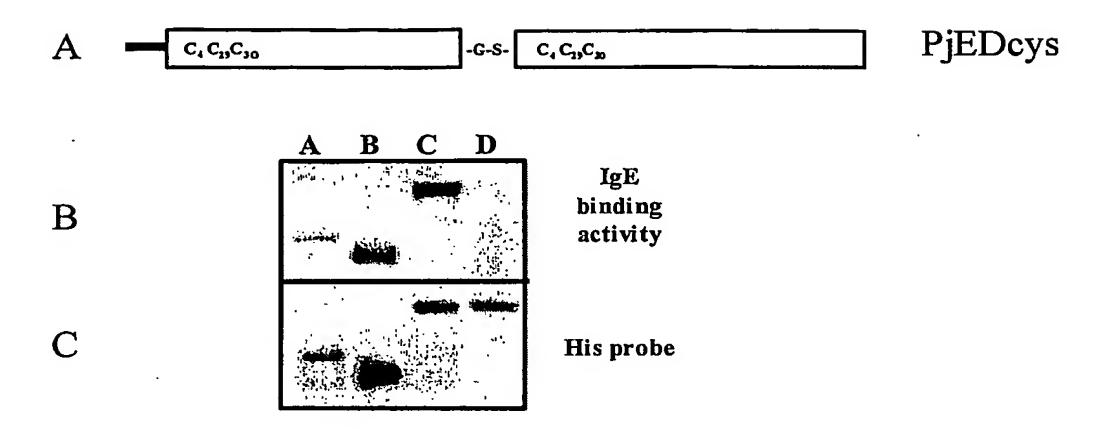


Fig 4

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Binding inhibition ELISA

	Dimer W.T	PjEDcys	
Patient 1	68%	7%	
Patient 2	56%	3,5%	
Patient 3	62%	10%	
Patient 4	62%	8%	
Patient 5	62%	8%	

Fig. 5

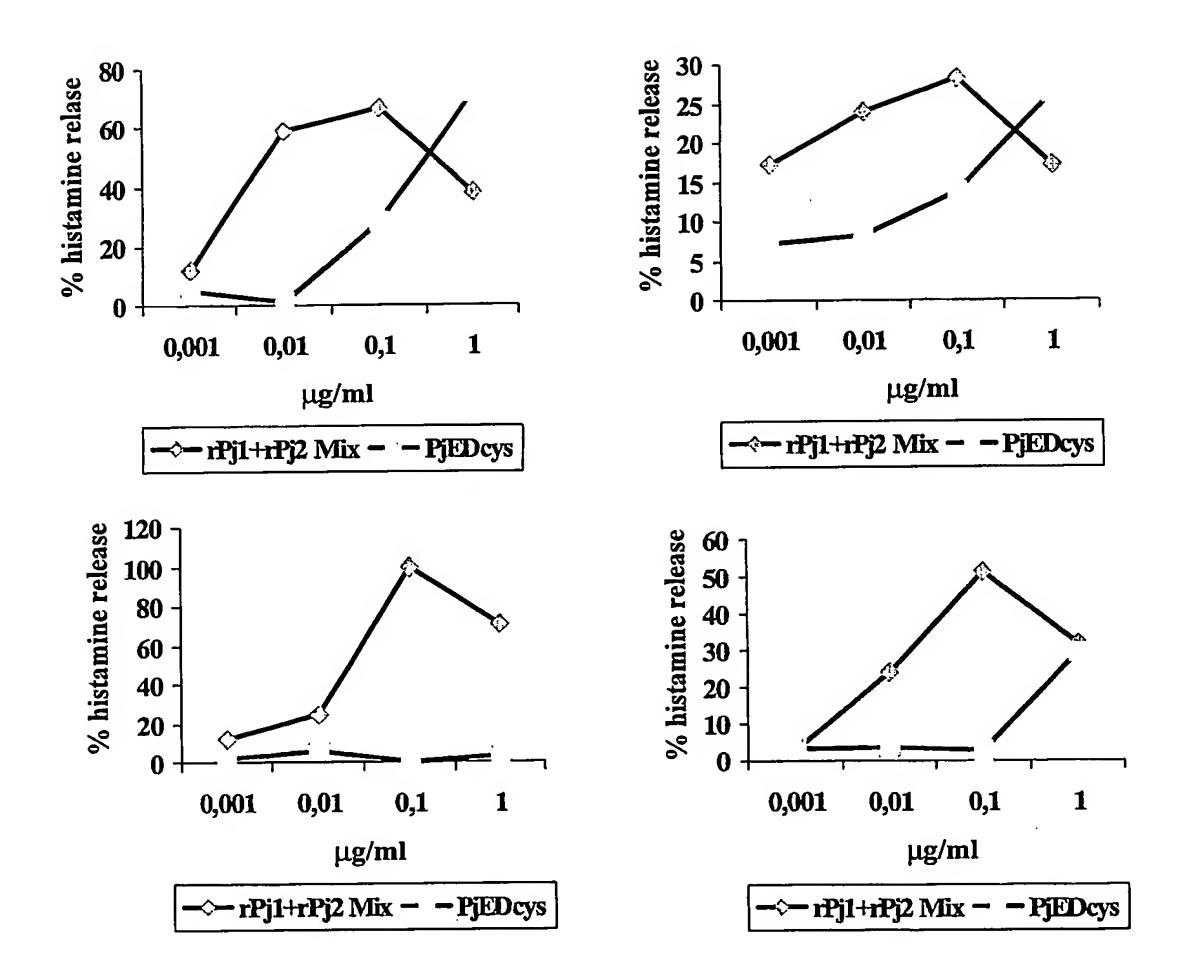
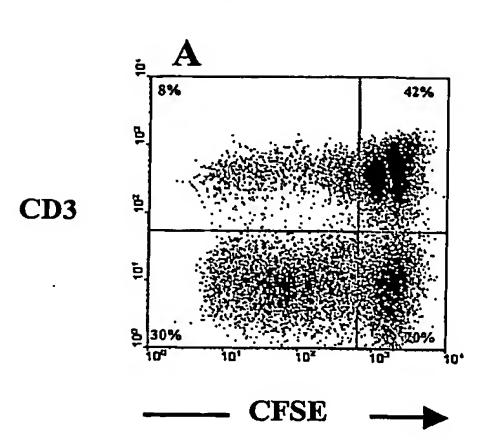
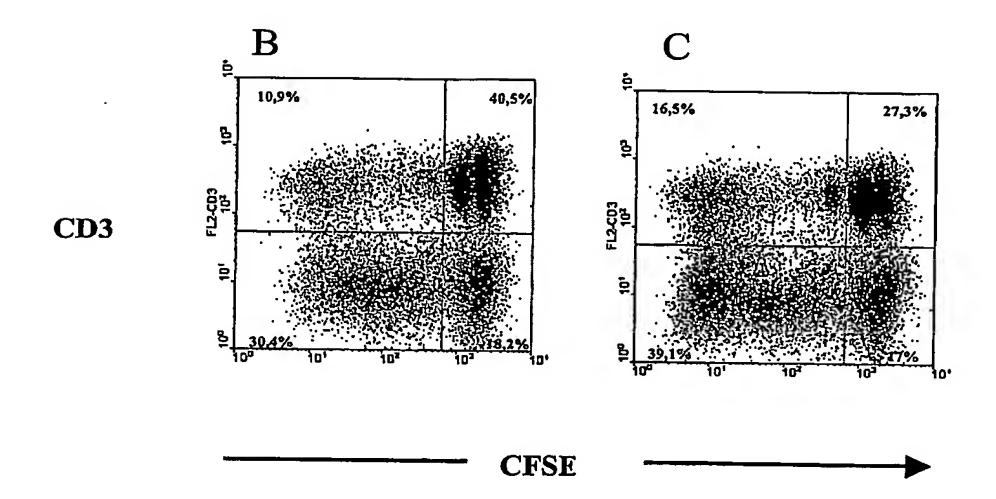


Fig. 6





		IVILX	PjEDcys
D		lµg/ml	lμg/ml
	Patient 1	2,9	8,5
	Patient 2	11.34	26.15
	Patient 3	3.3	3.62
	Patient 4	0.42	1.18
	Patient 5	10.44	18.86

Fig 7

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% of human IgE binding inhibition

P.judaica extract	ExPj	rParj1	rParj2	rParj1 rParj2
Patient 1	94	10	80	80
Patient 2	93	53	74	74
Patient 3	90	35	37	80
Patient 4	98	59	85	86
Patient 5	100	57	29	67
Patient 6	91	29	65	69
Patient 7	94	38	58	79
Average	94,3	40,1	61,1	76,5

Fig. 8

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% of human IgE binding inhibition

tested Protein	% of inhibition
rParj 1	95%
РjА	16%
PjB	85%
РjС	14%
PjD	15%

Fig. 9